

SEQUENCE LISTING

<110> The Curators of the University of Missouri

<120> LARGE SCALE EXPRESSION AND PURIFICATION OF RECOMBINANT
PROTEINS

<130> UMO1531.1

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<150> 60/218,125

<151> 2000-01-13

<160> 2

<170> PatentIn Ver. 2.1

<210> 1

<211> 4087

<212> DNA

<213> Bos taurus

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<221> CDS

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<222> (268)..(363)

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<222> (3178)

<223> A Poly (H) affinity tag comprising 6 His residues
have been inserted at the C-Terminus end of the
coding region of the protein

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ggcgaacatc aactcgtgct tgaaaaatac caacttggag cccggtttga gaagctacat 180

cagagtctcg agatgcgacg ctacaatctg cagttttcac tagcttccca gtaggttggg 240

acagttggaa ctctgccatt gcccagc atg ctg cag ttc agt ctg tca ccc acc 294

Met Leu Gln Phe Ser Leu Ser Pro Thr

1

5

ttg tcg atg gga ttt cac gtg ata gcc atg gtg gct ctc ttg ttt tcc 342

Leu Ser Met Gly Phe His Val Ile Ala Met Val Ala Leu Leu Phe Ser

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25

cat gtg gac cat ata agt gct gag aca gaa atg gaa gga gaa ggc aac 390

His Val Asp His Ile Ser Ala Glu Thr Glu Met Glu Gly Glu Gly Asn

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35

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gag act ggc gag tgt act ggc tcc tat tac tgt aag aag ggg gtg att 438

Glu Thr Gly Glu Cys Thr Gly Ser Tyr Tyr Cys Lys Lys Gly Val Ile

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tta ccc att tgg gag ccc cag gac cct tcc ttt gga gac aaa att gct 486

Leu Pro Ile Trp Glu Pro Gln Asp Pro Ser Phe Gly Asp Lys Ile Ala

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aga gcg act gtg tat ttt gtg gcc atg gtc tac atg ttt ctt gga gtc 534

Arg Ala Thr Val Tyr Phe Val Ala Met Val Tyr Met Phe Leu Gly Val

75

80

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tca atc att gct gac cgg ttc atg tcc tct ata gaa gtc atc acg tct 582

Ser Ile Ile Ala Asp Arg Phe Met Ser Ser Ile Glu Val Ile Thr Ser

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95

100

105

caa gag aaa gaa atc acc ata aag aaa ccc aat gga gag acc acc aag 630

Gln Glu Lys Glu Ile Thr Ile Lys Lys Pro Asn Gly Glu Thr Thr Lys

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aca act gtg agg atc tgg aat gag aca gtg tcc aac ctg acc ttg atg 678

Thr Thr Val Arg Ile Trp Asn Glu Thr Val Ser Asn Leu Thr Leu Met

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gcc ctg ggg tct tca gct cca gag att ctc ctt tca gta atc gag gtg 726

Ala Leu Gly Ser Ser Ala Pro Glu Ile Leu Leu Ser Val Ile Glu Val

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Cys Gly His Asn Phe Thr Ala Gly Asp Leu Gly Pro Ser Thr Ile Val

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ggg agt gct gca ttc aac atg ttc atc atc att gcc ctt tgt gtg tat 822

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180

185

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Gln Asp Asp Glu Glu Ala Arg Arg Glu Met Ala Arg Ile Leu Lys Glu	
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Leu Lys Gln Lys His Pro Glu Lys Glu Ile Glu Gln Leu Ile Glu Leu	
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Ala Asn Tyr Gln Val Leu Ser Gln Gln Gln Lys Ser Arg Ala Phe Tyr	
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cgt att caa gct acc cgc ctg atg acc gga gca ggc aac att tta aag	1398
Arg Ile Gln Ala Thr Arg Leu Met Thr Gly Ala Gly Asn Ile Leu Lys	
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Arg His Ala Ala Asp Gln Ala Arg Lys Ala Val Ser Met His Glu Val	
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Arg Thr Glu Asp Gly Thr Ala Asn Ala Gly Ser Asp Tyr Glu Phe Thr	
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Glu Gly Thr Val Val Phe Lys Pro Gly Glu Thr Gln Lys Glu Ile Arg	
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Val Gly Ile Ile Asp Asp Asp Ile Phe Glu Glu Asp Glu Asn Phe Leu	
475 480 485	
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Val His Leu Ser Asn Val Lys Val Ser Leu Glu Ala Ser Glu Asp Gly	
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Ile Leu Glu Ala Ser His Val Ser Thr Leu Ala Cys Leu Gly Ser Pro	
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Ser Thr Ala Thr Val Thr Ile Phe Asp Asp Asp His Ala Gly Ile Phe	
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<212> PRT

<213> Bos taurus

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Thr Ala Val Val	Phe Val Ala Leu Gly Thr Ser Val Pro Asp Thr Phe					
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Ala Ser Lys Val	Ala Ala Thr Gln Asp Gln Tyr Ala Asp Ala Ser Ile					
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Gly Asn Val Thr	Gly Ser Asn Ala Val Asn Val Phe Leu Gly Ile Gly					
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Val Ala Trp Ser	Ile Ala Ala Ile Tyr His Ala Ala Asn Gly Glu Gln					
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Glu Ile Gly Gly	Glu Leu Gly Gly Pro Arg Thr Ala Lys Leu Leu Thr					
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Ser Cys Leu Phe	Val Leu Leu Trp Leu Leu Tyr Ile Phe Phe Ser Ser					
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